

SEQUENCE LISTING

<110> Adams, Sean H.
Goddard, Audrey D.
Grimaldi, J. Christopher

<120> BFIT Compositions and Methods of Use

<130> 10716-3

<140> Unknown

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<170> PatentIn Ver. 2.1

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<211> 1857

<212> DNA

<213> Homo sapiens

<400> 1

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<211> 607

<212> PRT

<213> Homo sapiens

<400> 2

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Ser Ala Met Ala Asp Gly Glu Gly Tyr Arg Asn Pro Thr Glu Val Gln
 35 40 45

Met Ser Gln Leu Val Leu Pro Cys His Thr Asn Gln Arg Gly Glu Leu
 50 55 60

Ser Val Gly Gln Leu Leu Lys Trp Ile Asp Thr Thr Ala Cys Leu Ser
 65 70 75 80

Ala Glu Arg His Ala Gly Cys Pro Cys Val Thr Ala Ser Met Asp Asp
 85 90 95

Ile Tyr Phe Glu His Thr Ile Ser Val Gly Gln Val Val Asn Ile Lys
 100 105 110

Ala Lys Val Asn Arg Ala Phe Asn Ser Ser Met Glu Val Gly Ile Gln
 115 120 125

Val Ala Ser Glu Asp Leu Cys Ser Glu Lys Gln Trp Asn Val Cys Lys
 130 135 140

Ala Leu Ala Thr Phe Val Ala Arg Arg Glu Ile Thr Lys Val Lys Leu
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Lys Gln Ile Thr Pro Arg Thr Glu Glu Glu Lys Met Glu His Ser Val
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Ala Ala Glu Arg Arg Arg Met Arg Leu Val Tyr Ala Asp Thr Ile Lys

180

185

190

Asp Leu Leu Ala Asn Cys Ala Ile Gln Gly Asp Leu Glu Ser Arg Asp
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Cys Ser Arg Met Val Pro Ala Glu Lys Thr Arg Val Glu Ser Val Glu
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Leu Val Leu Pro Pro His Ala Asn His Gln Gly Asn Thr Phe Gly Gly
 225 230 235 240

Gln Ile Met Ala Trp Met Glu Asn Val Ala Thr Ile Ala Ala Ser Arg
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Leu Cys Arg Ala His Pro Thr Leu Lys Ala Ile Glu Met Phe His Phe
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Arg Gly Pro Ser Gln Val Gly Asp Arg Leu Val Leu Lys Ala Ile Val
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Asn Asn Ala Phe Lys His Ser Met Glu Val Gly Val Cys Val Glu Ala
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Tyr Arg Gln Glu Ala Glu Thr His Arg Arg His Ile Asn Ser Ala Phe
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Met Thr Phe Val Val Leu Asp Ala Asp Asp Gln Pro Gln Leu Leu Pro
 325 330 335

Trp Ile Arg Pro Gln Pro Gly Asp Gly Glu Arg Arg Tyr Arg Glu Ala
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Ser Ala Arg Lys Lys Ile Arg Leu Asp Arg Lys Tyr Ile Val Ser Cys
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Lys Gln Thr Glu Val Pro Leu Ser Val Pro Trp Asp Pro Ser Asn Gln
 370 375 380

Val Tyr Leu Ser Tyr Asn Asn Val Ser Ser Leu Lys Met Leu Val Ala
 385 390 395 400

Lys Asp Asn Trp Val Leu Ser Ser Glu Ile Ser Gln Val Arg Leu Tyr
 405 410 415

Thr Leu Glu Asp Asp Lys Phe Leu Ser Phe His Met Glu Met Val Val
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His Val Asp Ala Ala Gln Ala Phe Leu Leu Leu Ser Asp Leu Arg Gln

435

440

445

Arg Pro Glu Trp Asp Lys His Tyr Arg Ser Val Glu Leu Val Gln Gln
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Val Asp Glu Asp Asp Ala Ile Tyr His Val Thr Ser Pro Ala Leu Gly
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Gly His Thr Lys Pro Gln Asp Phe Val Ile Leu Ala Ser Arg Arg Lys
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Pro Cys Asp Asn Gly Asp Pro Tyr Val Ile Ala Leu Arg Ser Val Thr
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Leu Pro Thr His Arg Glu Thr Pro Glu Tyr Arg Arg Gly Glu Thr Leu
515 520 525

Cys Ser Gly Phe Cys Leu Trp Arg Glu Gly Asp Gln Leu Thr Lys Cys
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Cys Trp Val Arg Val Ser Leu Thr Glu Leu Val Ser Ala Ser Gly Phe
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Tyr Ser Trp Gly Leu Glu Ser Arg Ser Lys Gly Arg Arg Ser Asp Gly
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<210> 3

<211> 1818

<212> DNA

<213> Homo sapiens

<400> 3

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<210> 4

<211> 594

<212> PRT

<213> Homo sapiens

<400> 4

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Phe Ser Asn Arg Thr Ser Arg Lys Ser Ala Leu Arg Ala Gly Asn Asp

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25

30

Ser Ala Met Ala Asp Gly Glu Gly Tyr Arg Asn Pro Thr Glu Val Gln

35

40

45

Met Ser Gln Leu Val Leu Pro Cys His Thr Asn Gln Arg Gly Glu Leu

50

55

60

Ser Val Gly Gln Leu Leu Lys Trp Ile Asp Thr Thr Ala Cys Leu Ser

65

70

75

80

Ala Glu Arg His Ala Gly Cys Pro Cys Val Thr Ala Ser Met Asp Asp

85

90

95

Ile Tyr Phe Glu His Thr Ile Ser Val Gly Gln Val Val Asn Ile Lys			
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Ala Lys Val Asn Arg Ala Phe Asn Ser Ser Met Glu Val Gly Ile Gln			
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Val Ala Ser Glu Asp Leu Cys Ser Glu Lys Gln Trp Asn Val Cys Lys			
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Ala Leu Ala Thr Phe Val Ala Arg Arg Glu Ile Thr Lys Val Lys Leu			
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Lys Gln Ile Thr Pro Arg Thr Glu Glu Glu Lys Met Glu His Ser Val			
165	170	175	
Ala Ala Glu Arg Arg Arg Met Arg Leu Val Tyr Ala Asp Thr Ile Lys			
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Asp Leu Leu Ala Asn Cys Ala Ile Gln Gly Asp Leu Glu Ser Arg Asp			
195	200	205	
Cys Ser Arg Met Val Pro Ala Glu Lys Thr Arg Val Glu Ser Val Glu			
210	215	220	
Leu Val Leu Pro Pro His Ala Asn His Gln Gly Asn Thr Phe Gly Gly			
225	230	235	240
Gln Ile Met Ala Trp Met Glu Asn Val Ala Thr Ile Ala Ala Ser Arg			
245	250	255	
Leu Cys Arg Ala His Pro Thr Leu Lys Ala Ile Glu Met Phe His Phe			
260	265	270	
Arg Gly Pro Ser Gln Val Gly Asp Arg Leu Val Leu Lys Ala Ile Val			
275	280	285	
Asn Asn Ala Phe Lys His Ser Met Glu Val Gly Val Cys Val Glu Ala			
290	295	300	
Tyr Arg Gln Glu Ala Glu Thr His Arg Arg His Ile Asn Ser Ala Phe			
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Met Thr Phe Val Val Leu Asp Ala Asp Asp Gln Pro Gln Leu Leu Pro			
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Trp Ile Arg Pro Gln Pro Gly Asp Gly Glu Arg Arg Tyr Arg Glu Ala			
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Ser Ala Arg Lys Lys Ile Arg Leu Asp Arg Lys Tyr Ile Val Ser Cys
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Lys Gln Thr Glu Val Pro Leu Ser Val Pro Trp Asp Pro Ser Asn Gln
370 375 380

Val Tyr Leu Ser Tyr Asn Asn Val Ser Ser Leu Lys Met Leu Val Ala
385 390 395 400

Lys Asp Asn Trp Val Leu Ser Ser Glu Ile Ser Gln Val Arg Leu Tyr
405 410 415

Thr Leu Glu Asp Asp Lys Phe Leu Ser Phe His Met Glu Met Val Val
420 425 430

His Val Asp Ala Ala Gln Ala Phe Leu Leu Leu Ser Asp Leu Arg Gln
435 440 445

Arg Pro Glu Trp Asp Lys His Tyr Arg Ser Val Glu Leu Val Gln Gln
450 455 460

Val Asp Glu Asp Asp Ala Ile Tyr His Val Thr Ser Pro Ala Leu Gly
465 470 475 480

Gly His Thr Lys Pro Gln Asp Phe Val Ile Leu Ala Ser Arg Arg Lys
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Pro Cys Asp Asn Gly Asp Pro Tyr Val Ile Ala Leu Arg Ser Val Thr
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Leu Pro Thr His Arg Glu Thr Pro Glu Tyr Arg Arg Gly Glu Thr Leu
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Cys Ser Gly Phe Cys Leu Trp Arg Glu Gly Asp Gln Leu Thr Lys Val
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Ser Tyr Tyr Asn Gln Ala Thr Pro Gly Val Leu Asn Tyr Val Thr Thr
545 550 555 560

Asn Val Ala Gly Leu Ser Ser Glu Phe Tyr Thr Thr Phe Lys Ala Cys
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Thr Leu

<210> 5
 <211> 2699
 <212> DNA
 <213> Mus musculus

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<210> 6

<211> 594

<212> PRT

<213> Mus musculus

<400> 6

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Asp	Pro	Pro	Thr	Met	Ala	Glu	Gly	Glu	Gly	Tyr	Arg	Asn	Pro	Thr	Glu
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Val	Gln	Met	Ser	Gln	Leu	Val	Leu	Pro	Cys	His	Thr	Asn	His	Arg	Gly
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Glu	Leu	Ser	Ile	Gly	Gln	Leu	Leu	Lys	Trp	Ile	Asp	Thr	Thr	Ala	Cys
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Leu	Ser	Ala	Glu	Arg	His	Ala	Gly	Cys	Pro	Cys	Val	Thr	Ala	Ser	Met
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Asp	Asp	Ile	Tyr	Phe	Asp	His	Thr	Ile	Ser	Val	Gly	Gln	Val	Val	Asn
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Cys	Lys	Ala	Leu	Ala	Thr	Phe	Val	Ala	His	Arg	Glu	Leu	Ser	Lys	Val
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Lys	Leu	Lys	Gln	Val	Ile	Pro	Leu	Thr	Glu	Glu	Glu	Lys	Thr	Glu	His
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Gly	Val	Ala	Ala	Glu	Arg	Arg	Arg	Met	Arg	Leu	Val	Tyr	Ala	Asp	Thr
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Ile Lys Asp Leu Leu Thr His Cys Val Ile Gln Asp Asp Leu Asp Lys	195	200	205
Asp Cys Ser Asn Met Val Pro Ala Glu Lys Thr Arg Val Glu Ser Val	210	215	220
Glu Leu Val Leu Pro Pro His Ala Asn His Gln Gly Asn Thr Phe Gly	225	230	235 240
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Arg Leu Cys His Ala His Pro Thr Leu Lys Ala Ile Glu Met Phe His	260	265	270
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Val Asn Asn Ala Phe Lys His Ser Met Glu Val Gly Val Cys Val Glu	290	295	300
Ala Tyr Arg Gln Glu Ala Glu Thr Gln Arg Arg His Ile Asn Ser Ala	305	310	315 320
Phe Met Thr Phe Val Val Leu Asp Lys Asp Asp Gln Pro Gln Lys Leu	325	330	335
Pro Trp Ile Arg Pro Gln Pro Gly Glu Gly Glu Arg Arg Tyr Arg Glu	340	345	350
Ala Ser Ala Arg Lys Lys Ile Arg Leu Asp Arg Lys Tyr Leu Val Ser	355	360	365
Cys Lys Gln Ala Glu Val Ala Leu Ser Val Pro Trp Asp Pro Ser Asn	370	375	380
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Ala Lys Asp Asn Trp Val Leu Ser Val Glu Ile Ser Glu Val Arg Leu	405	410	415
Tyr Ile Leu Glu Glu Asp Phe Leu Ser Phe His Leu Glu Met Val Val	420	425	430
Asn Val Asp Ala Ala Gln Val Phe Gln Leu Leu Ser Asp Leu Arg Arg	435	440	445

Arg Pro Glu Trp Asp Lys His Tyr Arg Ser Val Glu Leu Val Gln Gln
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Val Asp Glu Asp Asp Ala Ile Tyr His Val Ile Ser Pro Ala Leu Ser
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Gly Asn Thr Lys Pro Gln Asp Phe Val Ile Leu Ala Ser Arg Arg Lys
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Pro Cys Asp Asn Gly Asp Pro Tyr Val Ile Ala Leu Arg Ser Val Thr
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Cys Ser Gly Phe Cys Leu Trp Arg Glu Gly Asp Gln Met Thr Lys Val
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Thr Leu

<210> 7

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Mouse BFIT
forward primer

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<210> 8

<211> 23

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Mouse BFIT
probe

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<210> 9
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<220>
<223> Description of Artificial Sequence: Mouse BFIT
reverse primer

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<210> 10
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<223> Description of Artificial Sequence: Human BFIT1
forward primer

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<210> 11
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<223> Description of Artificial Sequence: Human BFIT1
probe

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<223> Description of Artificial Sequence: Human BFIT1
reverse primer

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<210> 13
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<210> 14
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<220>
<223> Description of Artificial Sequence: Human BFIT2
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<210> 15
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